



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial-Number: 10/068,137A
Source: 1600
Date Processed by STIC: 4/5/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 2023 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

ERROR DETECTED **SUGGESTED CORRECTION** SERIAL NUMBER: 10/068,137A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics The number/text at the end of each line "wrapped" down to the next line. This may occur if your file
 Wrapped Aminos was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will
 prevent "wrapping."

- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

- 3 Misaligned Amino The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers;
 Numbering use space characters; instead.

- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please
 ensure your subsequent submission is saved in ASCII text.

- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. **Per Sequence Rules,**
 each n or Xaa can only represent a single residue. Please present the maximum number of each
 residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 PatentIn 2.0 A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid
 "bug" sequences(s) . Normally, PatentIn would automatically generate this section from the
 previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to
 the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for**
 Artificial or Unknown sequences.

- 7 Skipped Sequences Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (OLD RULES) (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

- 8 Skipped Sequences Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 (NEW RULES) <210> sequence id number
 <400> sequence id number
 000

- 9 Use of n's or Xaa's Use of n's and/or Xaa's have been detected in the Sequence Listing.
 (NEW RULES) Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

- 10 ✓ Invalid <213> Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or
 Response scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or
 is Artificial Sequence

- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or
 "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

- 12 PatentIn 2.0 Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file,
 "bug" resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence
 listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



1600

RAW SEQUENCE LISTING

DATE: 04/05/2004

PATENT APPLICATION: US/10/068,137A

TIME: 15:34:37

Input Set : A:\Sequence Listing 2-17-04 (copy to Examiner).txt

Output Set: N:\CRF4\04052004\J068137A.raw

2 <110> APPLICANT: Biopolo s.c.r.l.
 3 <120> TITLE OF INVENTION: YEAST STRAINS FOR THE PRODUCTION OF LACTIC ACID
 4 <130> FILE REFERENCE: 2027.547000

C--> 5 <140> CURRENT APPLICATION NUMBER: US/10/068,137A

C--> 6 <141> CURRENT FILING DATE: 2002-02-06

7 <150> PRIOR APPLICATION NUMBER: ITALY MI97A002080

8 <151> PRIOR FILING DATE: 1997-12-09

9 <150> PRIOR APPLICATION NUMBER: PCT/EP98/05758

10 <151> PRIOR FILING DATE: 1998-09-11

11 <160> NUMBER OF SEQ ID NOS: 12

12 <170> SOFTWARE: PatentIn version 3.0

13 <210> SEQ ID NO: 1

14 <211> LENGTH: 27

15 <212> TYPE: DNA

C--> 16 <213> ORGANISM: Artificial/Unknown

17 <220> FEATURE:

18 <221> NAME/KEY: misc_feature

19 <222> LOCATION: ()..()

20 <223> OTHER INFORMATION: Oligomer anneals to inactivated ampicillin resistance gene of
 21 plasmid pALTER-land restores ampicillin resistance to mutant
 22 strand after mutagenesis reaction

24 <400> SEQUENCE: 1

25 gttgccattg ctgcaggcat cgtggtg

29 <210> SEQ ID NO: 2

30 <211> LENGTH: 29

31 <212> TYPE: DNA

C--> 32 <213> ORGANISM: Artificial/Unknown

33 <220> FEATURE:

34 <221> NAME/KEY: misc_feature

35 <222> LOCATION: ()..()

36 <223> OTHER INFORMATION: Oligomer anneals upstream of Bos taurus LDH gene in 5'UTR and
 37 creates XbaI restriction site 11 bp before start codon after
 38 mutagenesis reaction

39 <400> SEQUENCE: 2

40 ccttttaggt ctagatccaa gatggcaac

44 <210> SEQ ID NO: 3

45 <211> LENGTH: 27

46 <212> TYPE: DNA

C--> 47 <213> ORGANISM: Artificial/Unknown

48 <220> FEATURE:

49 <221> NAME/KEY: misc_feature

50 <222> LOCATION: ()..()

51 <223> OTHER INFORMATION: Oligomer anneals to 5' end of L. casei LDH gene and

introduces a

pgs 1-4
 Does Not Comply
 Corrected Diskette Needed

response

(see item 10 on Error Summary
 sheet)

Insert a space between "pALTER-1" and "and"

27

29

RAW SEQUENCE LISTING

DATE: 04/05/2004

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TIME: 15:34:37

Input Set : A:\Sequence Listing 2-17-04 (copy to Examiner).txt

Output Set: N:\CRF4\04052004\J068137A.raw

```

52      NcoI restriction site which changes the GTG start codon to an ATG
53      start codon
55 <400> SEQUENCE: 3
56 ccatggcaag tattacggat aaggatc                27
60 <210> SEQ ID NO: 4
61 <211> LENGTH: 24
62 <212> TYPE: DNA
63 <213> ORGANISM: Lactobacillus casei
64 <220> FEATURE:
65 <221> NAME/KEY: misc_feature
66 <222> LOCATION: ()..()
67 <223> OTHER INFORMATION: Antisense oligomer anneals 8 bp upstream of 3' end of L.
casei
68      LDH gene
70 <400> SEQUENCE: 4
71 ctatcactgc aggtttcga tgtc                24
75 <210> SEQ ID NO: 5
76 <211> LENGTH: 59
77 <212> TYPE: DNA
C--> 78 <213> ORGANISM: Artificial/Unknown
79 <220> FEATURE:
80 <221> NAME/KEY: misc_feature
81 <222> LOCATION: ()..()
82 <223> OTHER INFORMATION: Oligomer complementary to S. cerevisiae PDC1 and anneals to
83      loxP -KanSRD-loxP cassette
85 <400> SEQUENCE: 5
86 ttctactcat aacctcacgc aaaataacac agtcaaatca cagctgaagc ttcgtacgc        59
90 <210> SEQ ID NO: 6
91 <211> LENGTH: 62
92 <212> TYPE: DNA
C--> 93 <213> ORGANISM: Artificial/Unknown
94 <220> FEATURE:
95 <221> NAME/KEY: misc_feature
96 <222> LOCATION: ()..()
97 <223> OTHER INFORMATION: Antisense oligomer complementary to S. cerevisiae PDC1 and
anneal
98      to loxP-KanSRD-loxP cassette
100 <400> SEQUENCE: 6
101 aatgcttata aaactttaac taataattag agattaaatc gcataggcca ctagtggatc        60
103 tg                62
107 <210> SEQ ID NO: 7
108 <211> LENGTH: 59
109 <212> TYPE: DNA
C--> 110 <213> ORGANISM: Artificial/Unknown
111 <220> FEATURE:
112 <221> NAME/KEY: misc_feature
113 <222> LOCATION: ()..()
114 <223> OTHER INFORMATION: Oligomer complementary to S. cerevisiae PDC5 and anneals to
115      loxP -KanSRD-loxP cassette
117 <400> SEQUENCE: 7
118 atcaatctca aagagaacaa cacaatacaa taacaagaag cagctgaagc ttcgtacgc        59

```

RAW SEQUENCE LISTING

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PATENT APPLICATION: US/10/068,137A

TIME: 15:34:37

Input Set : A:\Sequence Listing 2-17-04 (copy to Examiner).txt

Output Set: N:\CRF4\04052004\J068137A.raw

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122 <210> SEQ ID NO: 8
123 <211> LENGTH: 62
124 <212> TYPE: DNA
C--> 125 <213> ORGANISM: Artificial/Unknown
126 <220> FEATURE:
127 <221> NAME/KEY: misc_feature
128 <222> LOCATION: ()..()
129 <223> OTHER INFORMATION: Antisense oligomer complementary to S. cerevisiae PDC5 and
anneal to
130         to loxP-KanSRD-loxP cassette
132 <400> SEQUENCE: 8
133 aaaatacaca aacgttgaat catgagtttt atgttaatta gcataggcca ctagtggatc      60
135 tg                                                                    62
139 <210> SEQ ID NO: 9
140 <211> LENGTH: 59
141 <212> TYPE: DNA
C--> 142 <213> ORGANISM: Artificial/Unknown
143 <220> FEATURE:
144 <221> NAME/KEY: misc_feature
145 <222> LOCATION: ()..()
146 <223> OTHER INFORMATION: Oligomer complementary to S. cerevisiae PDC6 and anneals to
147         loxP-KanSRD-loxP cassette
149 <400> SEQUENCE: 9
150 taaataaaaa acccacgtaa tatagcaaaa acatattgcc cagctgaagc ttcgtacgc      59
154 <210> SEQ ID NO: 10
155 <211> LENGTH: 62
156 <212> TYPE: DNA
C--> 157 <213> ORGANISM: Artificial/Unknown
158 <220> FEATURE:
159 <221> NAME/KEY: misc_feature
160 <222> LOCATION: ()..()
161 <223> OTHER INFORMATION: Antisense oligomer complementary to S. cerevisiae PDC6 and
anneals
162         to loxP-KanSRD-loxP cassette
164 <400> SEQUENCE: 10
165 tttatttgca acaataattc gtttgagtac actactaatg gcataggcca ctagtggatc      60
167 tg                                                                    62
171 <210> SEQ ID NO: 11
172 <211> LENGTH: 59
173 <212> TYPE: DNA
C--> 174 <213> ORGANISM: Artificial/Unknown
175 <220> FEATURE:
176 <221> NAME/KEY: misc_feature
177 <222> LOCATION: ()..()
178 <223> OTHER INFORMATION: Oligomer complementary to S. cerevisiae PDC2 and anneals to
179         loxP-KanSRD-loxP cassette
181 <400> SEQUENCE: 11
182 acgcaacttg aattggcaaa atgggcttat gagacgttcc cagctgaagc ttcgtacgc      59
186 <210> SEQ ID NO: 12
187 <211> LENGTH: 62
188 <212> TYPE: DNA

```

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TIME: 15:34:37

Input Set : A:\Sequence Listing 2-17-04 (copy to Examiner).txt

Output Set: N:\CRF4\04052004\J068137A.raw

C--> 189 <213> ORGANISM: Artificial/Unknown

190 <220> FEATURE:

191 <221> NAME/KEY: misc_feature

192 <222> LOCATION: ()..()

193 <223> OTHER INFORMATION: Antisense oligomer complementary to S. cerevisiae PDC2 and
anneals

194 to loxP-KanSRD-loxP cassette

196 <400> SEQUENCE: 12

197 agcctgtgtt accaggtaag tgtaagttat tagagtctgg gcataggcca ctagtggatc 60

199 tg 62

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/068,137A

DATE: 04/05/2004

TIME: 15:34:38

Input Set : A:\Sequence Listing 2-17-04 (copy to Examiner).txt

Output Set: N:\CRF4\04052004\J068137A.raw

L:5 M:270 C: Current Application Number differs, Replaced Current Application Number
 L:6 M:271 C: Current Filing Date differs, Replaced Current Filing Date
 L:16 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
 L:32 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
 L:47 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
 L:78 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:5
 L:93 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:6
 L:110 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7
 L:125 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:8
 L:142 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:9
 L:157 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:10
 L:174 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:11
 L:189 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:12